



# Preclinical Development of LYL119, a ROR1-Targeted CAR T-Cell Product Incorporating Four Novel T-Cell Reprogramming Technologies to Overcome Barriers to Effective Cell Therapy for Solid Tumors

Abstract 278

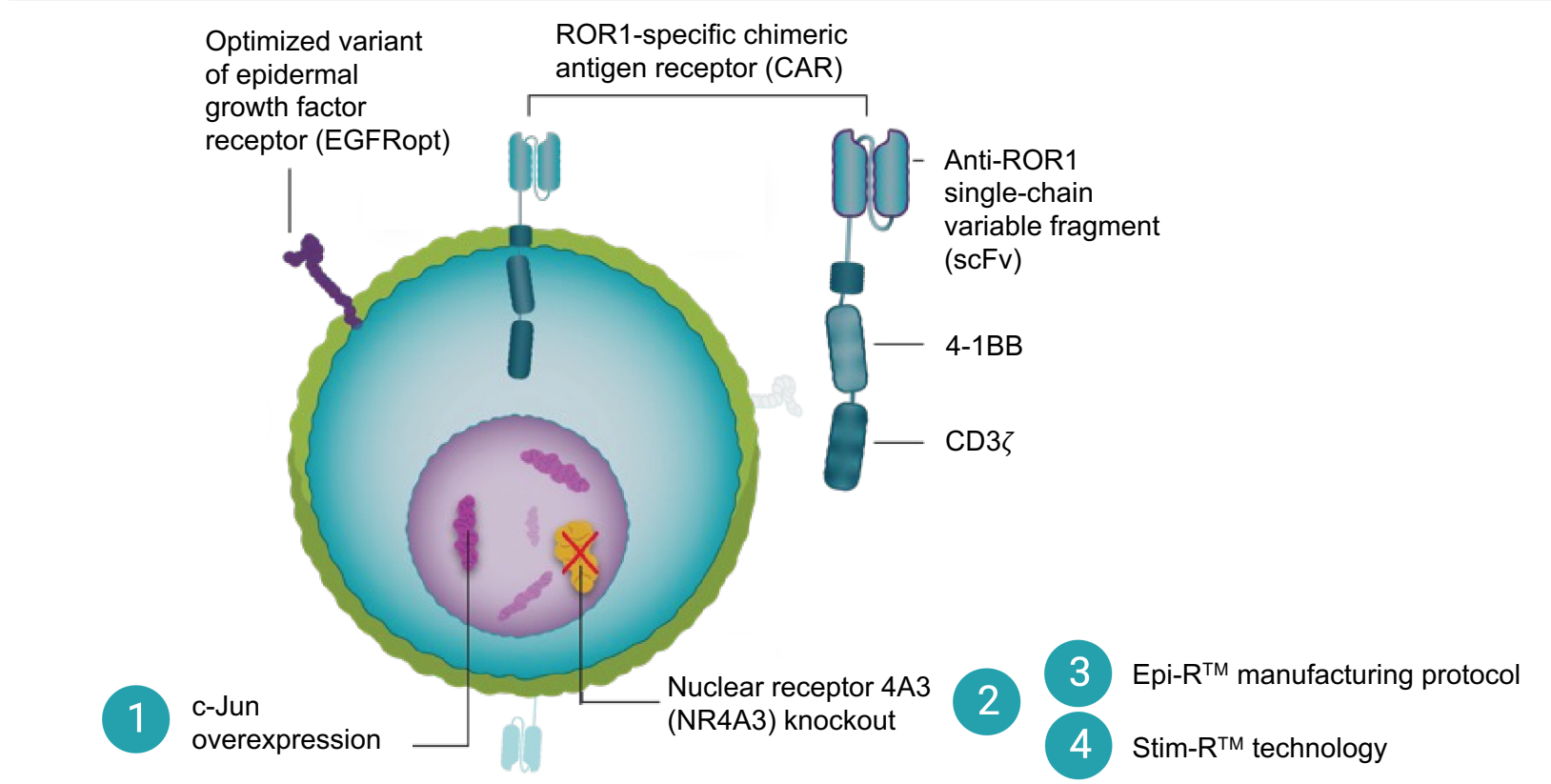
Viola C. Lam, Aileen Li, Meri Galindo Casas, Jessica Barragan, Christina Cheung, Jessica Briones, Esha Afreen, Grant Vavra, Jia Lu, Purnima Sundar, Rowena Martinez, Candace Sims, Shobha Potluri, Omar Ali, Alexander S. Cheung, and Rachel C. Lynn

Lyell Immunopharma, Inc., South San Francisco, CA and Seattle, WA

## Background

- T-cell exhaustion and lack of durable stemness (defined as the ability of cells to proliferate, persist, and self-renew) are key barriers to effective T-cell therapy in solid tumors<sup>1-2</sup>
- Lyell has developed multiple genetic and epigenetic T-cell reprogramming strategies to overcome these barriers:
  - Genetically reprogramming T cells through c-Jun overexpression delays exhaustion and results in increased proliferation, sustained cytokine production, and durable antitumor activity<sup>1,3-4</sup>
  - Genetically reprogramming T cells through *NR4A3* gene KO in combination with c-Jun overexpression further enhances resistance to exhaustion and improves antitumor activity<sup>5</sup>
  - Epigenetic reprogramming with Lyell's Epi-R™ manufacturing protocol preserves stem-like qualities by controlling T-cell proliferation and differentiation with optimized proprietary cell culture media and other manufacturing steps<sup>1,6-8</sup>
  - Epigenetic reprogramming with Lyell's Stim-R™ technology (a synthetic biomimetic designed to precisely and physiologically present T-cell activation signals during manufacturing) further improves T-cell polyfunctionality, persistence, and antitumor activity<sup>9</sup>
- These four T-cell reprogramming technologies are combined in LYL119, an investigational ROR1-targeted CAR T-cell product enhanced with c-Jun overexpression, *NR4A3* KO, and Epi-R and Stim-R technologies to overcome barriers to successful T-cell therapy in solid tumors (Figure 1)

Figure 1: LYL119, an investigational ROR1-targeted CAR T-cell product enhanced with Lyell's reprogramming technologies

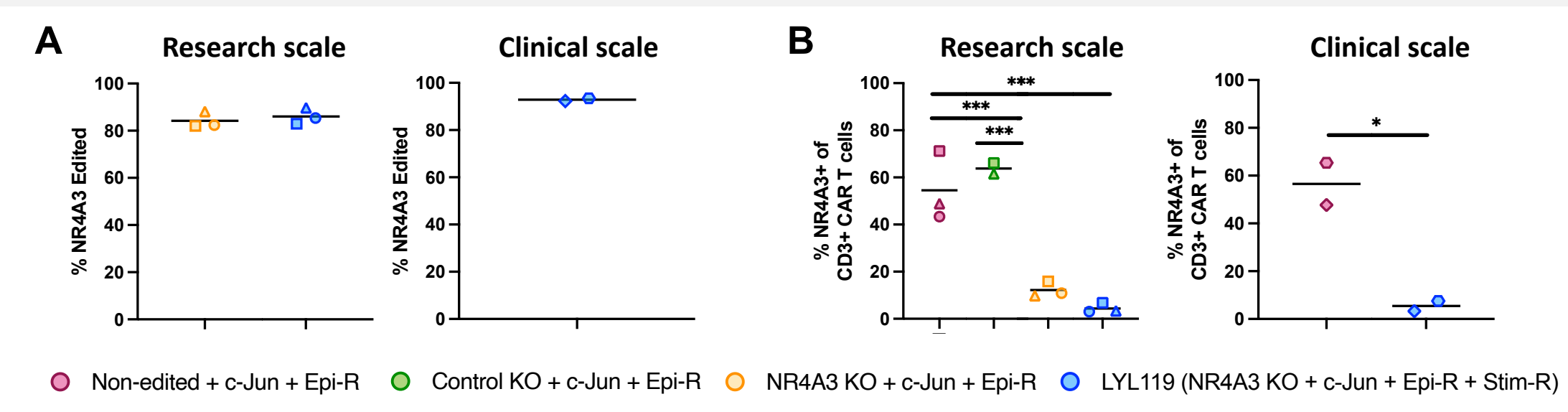


## Results

### High *NR4A3* genomic editing results in significantly reduced *NR4A3* protein expression

- NR4A3* editing efficiency was similar between *NR4A3* KO ROR1 CAR T cells manufactured with or without Stim-R technology and between research and clinical-scale products (Figure 2)
- NR4A3* protein reduction in LYL119 was similar between production scales (Figure 2)

Figure 2: LYL119 demonstrates high *NR4A3* editing resulting in reduced *NR4A3* protein expression

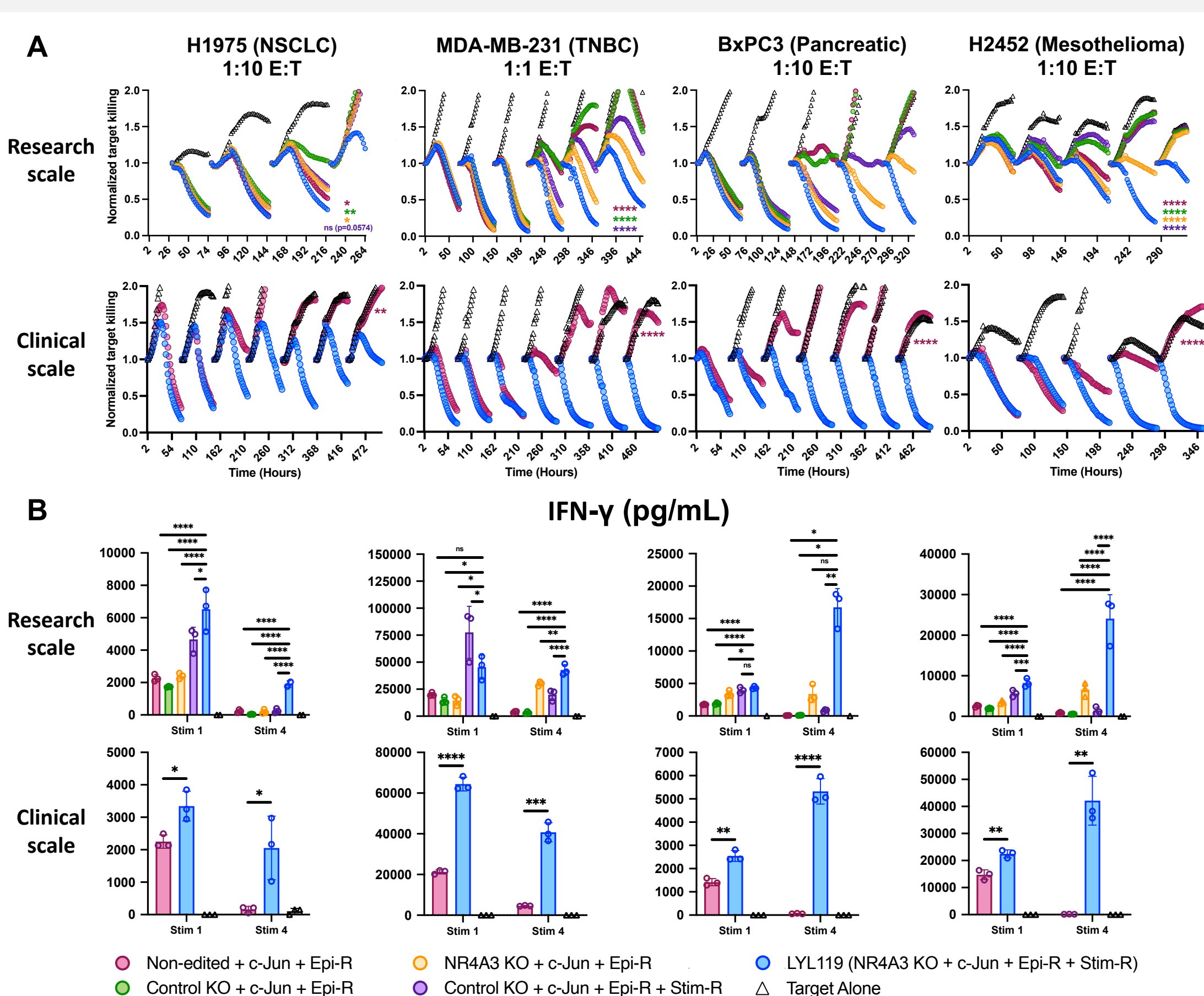


*NR4A3* genomic editing efficiency (A) and *NR4A3* protein expression (B) in edited ROR1 CAR T cells. Symbols correspond to ROR1 CAR T cells derived from independent donors (n=3 for research and n=2 for clinical scale). Asterisks indicate significant differences. \*p<0.05; \*\*\*p<0.001 by Tukey's ordinary one-way ANOVA statistical analysis (research) or an unpaired t-test (clinical).

### LYL119 demonstrates superior in vitro activity compared to ROR1 CAR T cells lacking one or more reprogramming technologies

Research and clinical-scale LYL119 demonstrated prolonged cytotoxicity and enhanced IFN- $\gamma$  secretion upon antigen restimulation with multiple different ROR1-expressing solid tumor cell lines (Figure 3)

Figure 3: LYL119 demonstrates superior cytotoxicity and enhanced IFN- $\gamma$  secretion



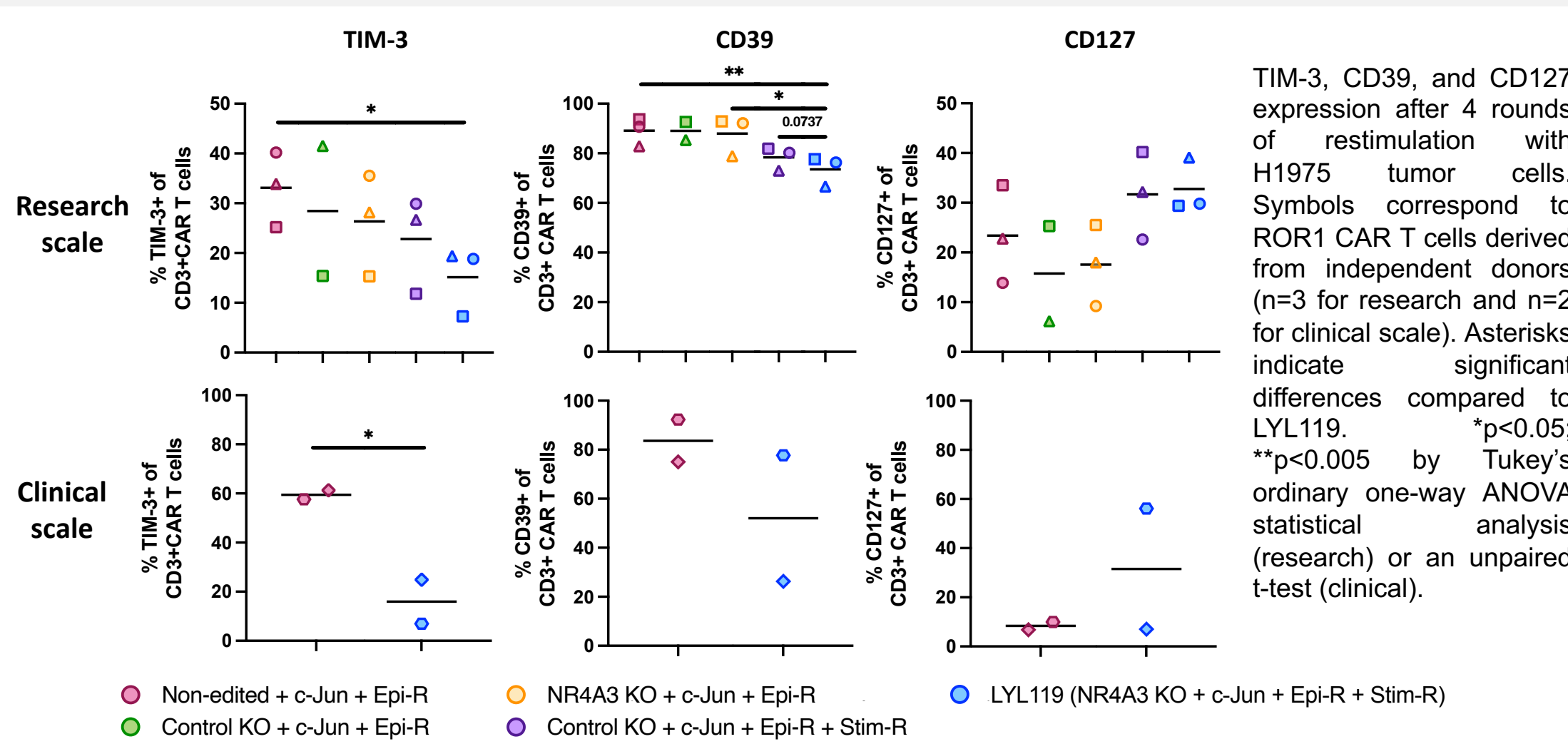
(A) Normalized target killing following sequential stimulation with ROR1<sup>+</sup> solid tumor cell lines. Lysis of NuLight Red (NLR)-expressing tumor cells was quantified by measuring total NLR intensity and normalized relative to the starting intensity for each round of stimulation. (B) IFN- $\gamma$  secretion during sequential exposure to antigen described in (A). Error bars represent mean  $\pm$  SD of triplicate wells. One representative donor is shown at research (n=3) and clinical (n=2) scale. Asterisks indicate significant differences compared to LYL119 (last timepoint of the last stimulation in A). ns-not significant, \*p<0.05; \*\*p<0.005; \*\*\*p<0.001; \*\*\*\*p<0.0001 by Tukey's ordinary one-way ANOVA statistical analysis.

### LYL119 exhibits an improved phenotype following antigen restimulation

Compared to non-edited ROR1 CAR T cells with c-Jun overexpression alone, LYL119 displayed:

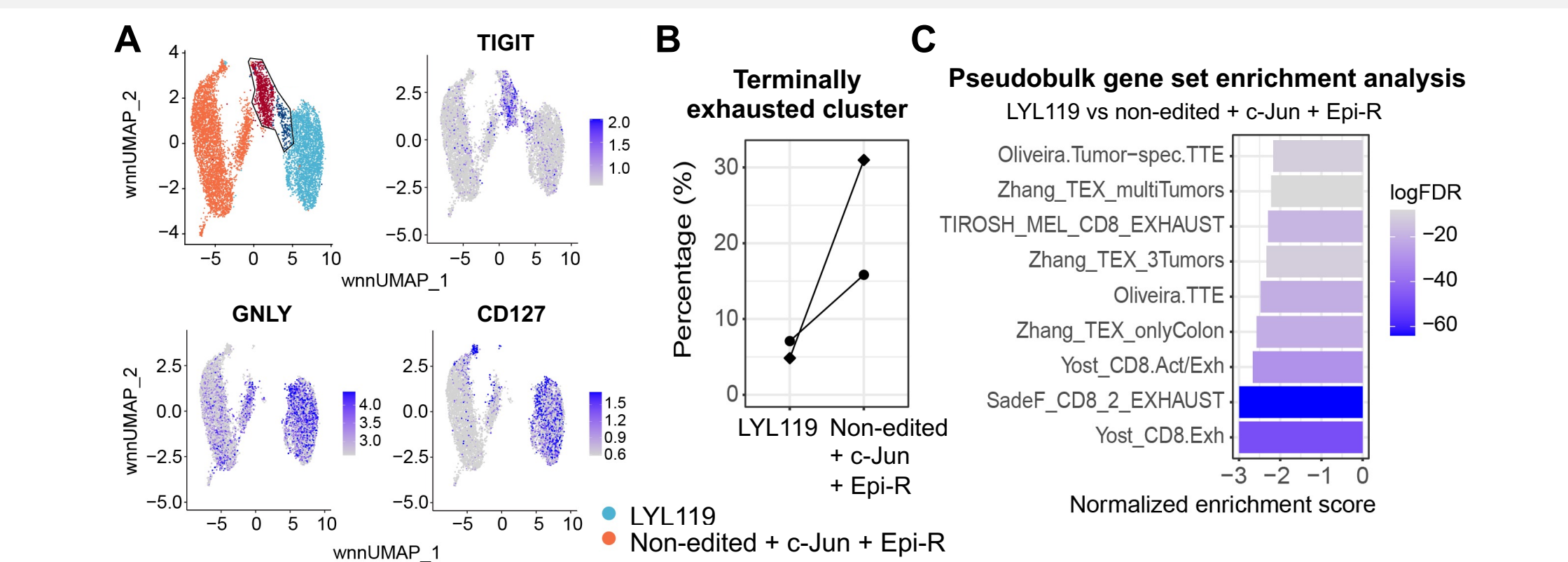
- Further reduction of T-cell exhaustion – Lower TIM-3 and CD39 expression (Figure 4), lower proportion of terminally exhausted cluster and significant down-regulation of multiple exhaustion-associated gene sets (Figure 5)
- Elevated CD127 expression (Figure 4, Figure 5), a marker for memory-like T cells, suggesting maintenance of stem-like characteristics
- Global up-regulation of GNLy, an effector-associated marker, suggesting enhanced effector-like characteristics (Figure 5)

Figure 4: LYL119 displays reduced T-cell exhaustion and enhanced memory-like phenotype following antigen restimulation



TIM-3, CD39, and CD127 expression after 4 rounds of restimulation with H1975 tumor cells. Symbols correspond to ROR1 CAR T cells derived from independent donors (n=3 for research and n=2 for clinical scale). Asterisks indicate significant differences compared to LYL119. \*p<0.05; \*\*p<0.005 by Tukey's ordinary one-way ANOVA statistical analysis (research) or an unpaired t-test (clinical).

Figure 5: LYL119 exhibits reduced T-cell exhaustion and enhanced memory-like and effector-like transcriptomic signatures after antigen restimulation



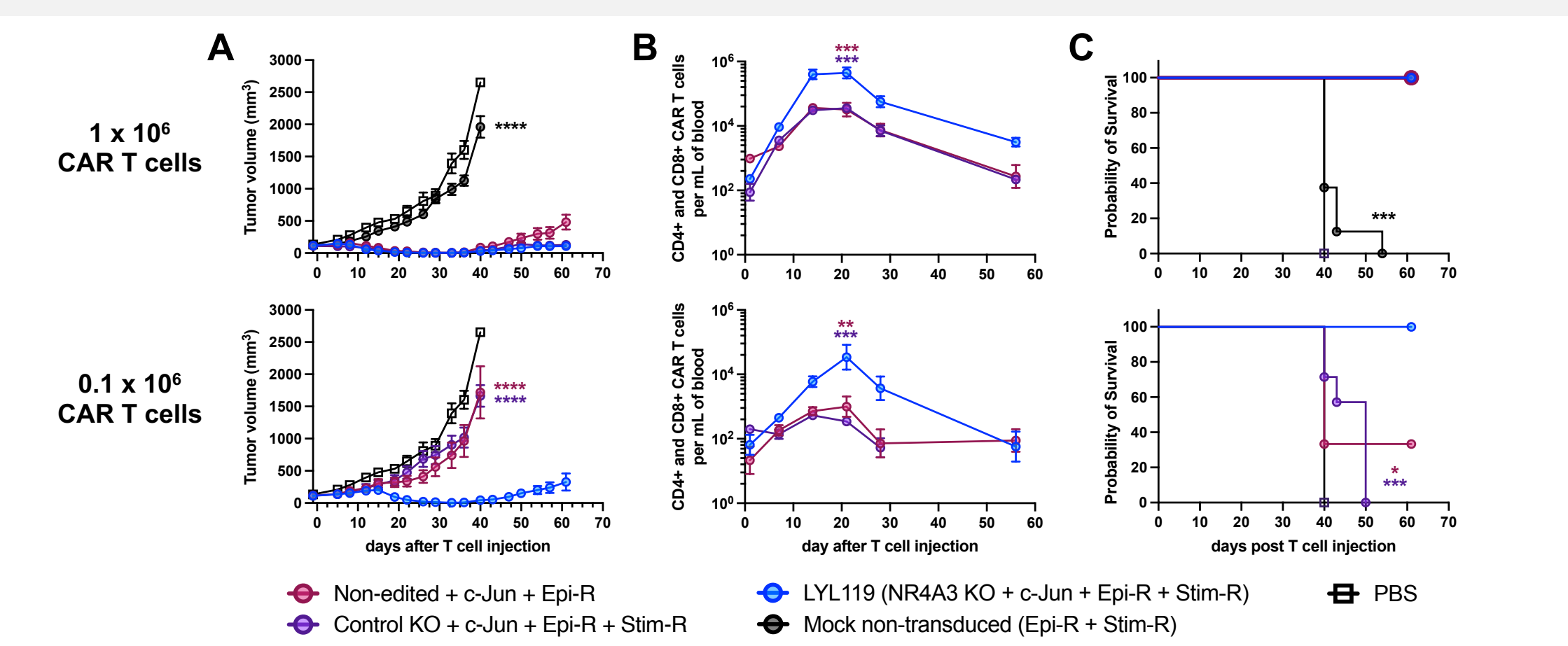
Single-cell Multiome data of LYL119 (blue) and non-edited + c-Jun + Epi-R (red) CD8<sup>+</sup> ROR1 CAR<sup>+</sup> T cells after 5 rounds of antigen restimulation with A549 tumor cells. (A) Representative UMAP plot derived from one clinical-scale donor (n=2) showing terminally exhausted clusters (circled and in dark blue for LYL119 and dark red for non-edited + c-Jun + Epi-R CAR T cells) enriched for TIGIT RNA expression. LYL119 shows enriched GNLy and CD127 RNA expressions. (B) Percentage of terminally exhausted cluster shown in A. Symbols correspond to CAR T cells derived from independent donors. (C) Exhaustion-associated gene set enrichment analysis.

### LYL119 has potent antitumor activity in vivo

Compared to non-edited ROR1 CAR T cells with c-Jun overexpression alone, LYL119:

- Demonstrated robust antitumor activity across a 10-fold CAR T-cell dose range
- Exhibited superior CAR T-cell expansion in the peripheral blood ranging 13- to 62-fold higher on Day 21 after T-cell injection
- Significantly improved animal survival at the low 0.1  $\times 10^6$  CAR T-cell dose (Figure 6)

Figure 6: LYL119 demonstrates robust antitumor activity in vivo

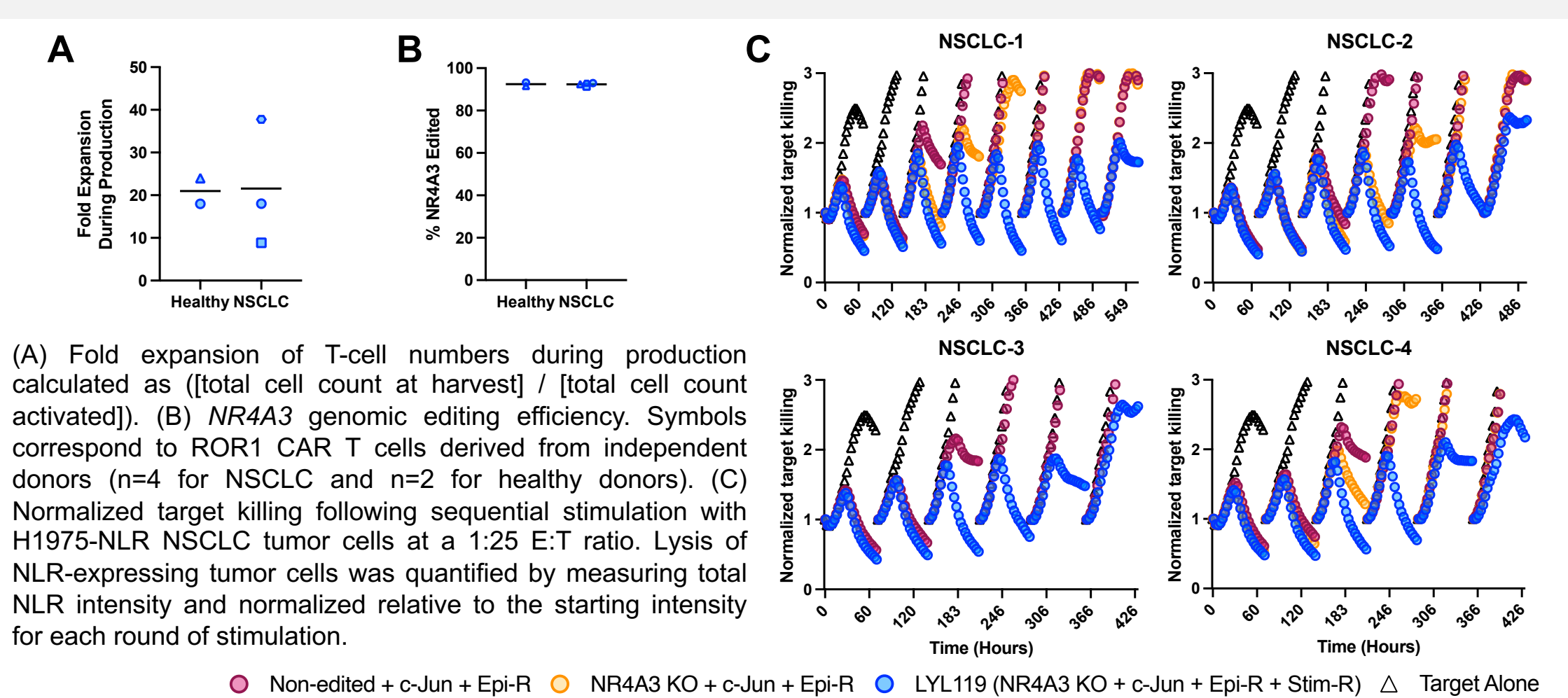


(A) Tumor volume, (B) peripheral blood ROR1 CAR T cells, and (C) animal survival at 1  $\times 10^6$  and 0.1  $\times 10^6$  CAR T-cell dose range in a H1975 xenograft NSG HLA dKO mouse model. Data from one representative research-scale donor of 3 independent animal studies (n=2 for research and n=1 for clinical scale) with n=6-10 mice/group is shown. Mean tumor volume curves were truncated at timepoints when less than 80% of animals/group remained on study. Error bars represent mean  $\pm$  SEM. Statistical analysis of peripheral blood ROR1 CAR T cell expansion was performed on Day 21 after T-cell injection. Asterisks indicate significant differences compared to LYL119-treated animals. \*p<0.05; \*\*p<0.005; \*\*\*p<0.001; \*\*\*\*p<0.0001 by Tukey's mixed-effects one-way ANOVA (A, B) or log-rank Mantel-Cox (C) statistical analyses.

### NSCLC patient-derived LYL119 products demonstrate robust cytotoxicity in vitro

- NSCLC patient-derived LYL119 demonstrated similar T-cell fold expansion as healthy donor-derived LYL119 during production
- Approximately 90% *NR4A3* genomic editing efficiency was achieved in NSCLC patient-derived LYL119 products
- NSCLC patient-derived LYL119 demonstrated superior cytotoxicity compared to patient ROR1 CAR T cells lacking 1 or 2 reprogramming technologies (Figure 7)

Figure 7: NSCLC patient-derived LYL119 demonstrates superior cytotoxicity in vitro



(A) Fold expansion of T-cell numbers during production calculated as ((total cell count at harvest) / [total cell count activated]). (B) *NR4A3* genomic editing efficiency. Symbols correspond to ROR1 CAR T cells derived from independent donors (n=4 for NSCLC and n=2 for healthy donors). (C) Normalized target killing following sequential stimulation with H1975-NLR NSCLC tumor cells at a 1:25 E:T ratio. Lysis of NLR-expressing tumor cells was quantified by measuring total NLR intensity and normalized relative to the starting intensity for each round of stimulation.

## Conclusions

- LYL119, which combines c-Jun overexpression, *NR4A3* KO, Epi-R protocol, and Stim-R technology exhibited:
  - Potent cytotoxicity and enhanced cytokine secretion upon antigen restimulation with multiple ROR1<sup>+</sup> solid tumor cell lines in vitro
  - An enhanced memory-like and effector-like phenotype with reduced T-cell exhaustion after antigen restimulation in vitro
  - Potent antitumor activity, superior ROR1 CAR T-cell expansion, and improved survival in a H1975 xenograft tumor mouse model
  - LYL119 produced from NSCLC patient cells demonstrated superior cytotoxic activity compared to ROR1 CAR T cells lacking 1 or 2 reprogramming technologies

These nonclinical studies further support the development of LYL119 for its potential to provide effective and durable CAR T-cell antitumor activity in patients with ROR1<sup>+</sup> solid tumors.

## Methods

- Healthy donor ROR1 CAR T cells were manufactured at research or clinical scale using the Epi-R protocol, activated with Stim-R technology or a standard reagent, and transduced with a lentiviral vector encoding c-Jun, the ROR1 CAR, and an optimized variant of EGFR (EGFRopt). NSCLC patient donor ROR1 CAR T cells were manufactured at research scale.
- T cells were electroporated with a single guide RNA targeting human *NR4A3* or control *CD19* (not expressed in T cells) complexed with SpyFl™ Cas9 nuclease (Aldevron®)
- ROR1 CAR T-cell cytotoxicity, cytokine production, and phenotype were evaluated in vitro following antigen restimulation assays designed to promote T-cell exhaustion
- Antitumor activity of ROR1 CAR T cells was evaluated in vivo using a ROR1-expressing H1975 human NSCLC xenograft model in NSG HLA dKO mice



## Abbreviations

4-1BB, tumor necrosis factor ligand superfamily member 9; CAR, chimeric antigen receptor; CD, cluster of differentiation; dKO, double knockout; EGFRopt, optimized variant of epidermal growth factor receptor; E:T, effector-to-target; GNLy, granulysin; HLA, human leukocyte antigen; IFN- $\gamma$ , interferon gamma; IL-2, interleukin 2; KO, knockout; NR4A3, nuclear receptor subfamily 4 group A member 3; NLR, NuLight Red; NSCLC, non-small cell lung cancer; NSG, NOD scid gamma; ROR1, receptor tyrosine kinase-like orphan receptor 1; scFv, single-chain variable fragment; SD, standard deviation; SEM, standard error of the mean; TIGIT, T cell immunoreceptor with Ig and ITIM domains; TIM-3, T cell immunoglobulin and mucin domain-containing protein 3; TNBC, triple negative breast cancer

## References

- Park S, et al. AACR 2022 Poster. Abstract 2754, 2. Krishna S, et al. Science. 2020;370:1328–1334, 3. Lynn RC, et al. Nature. 2019;576:293–300, 4. Park S, et al. ASGCT 2022 Poster. Abstract 661, 5. Lam V, et al. SITC 2022 Poster. Abstract 243, 6. Patel Y, et al. SITC 2022 Poster. Abstract 370, 7. Harris BD, et al. SITC 2022 Poster. Abstract 340, 8. Patel Y, et al. AACR Special Conference: Tumor Immunology and Immunotherapy 2022 Poster. Abstract A54, 9. Li A, et al. SITC 2022 Poster. Abstract 252.

## Acknowledgments

We thank members of Lyell Immunopharma's In Vivo Pharmacology team (Shawn Anderson, Ally Torres Diaz, Eskedar Nigatu, Christina Ta, and Nik Vad), NGS/Flow team (Sahithi Cheemalamarri, Emily Fu-Sum, Smita Ghanekar, Sheila Lou, Taran Ramadoss, Stefan Siebert, Ken Xiong, and Lora Zhao), Research team (Spencer Park, Sydney Spadinger, and Chiara Victor), and Vector Sciences team (David Anderson and Martin Wohlfahrt) for their experimental contributions.

Presented at SITC Annual Meeting 2023; Nov 1–5; San Diego, CA, USA